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<input type="checkbox"/>	L7	(vitamin adj B6) same L1	27
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<input type="checkbox"/>	L2	(gene or sequence or polynucleotide) same L1	251
<input type="checkbox"/>	L1	((erythronate-4-phosphate adj dehydrogenase)or (4-phosphoerythronate adj dehydrogenase)or (phosphoerythronate adj dehydrogenase) or pdx?)	815

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# NiceZyme View of ENZYME: EC 1.1.1.290

## Official Name

**4-phosphoerythronate dehydrogenase.**

## Alternative Name(s)

**4-O-phosphoerythronate dehydrogenase.**

**4PE dehydrogenase.**

**Erythronate-4-phosphate dehydrogenase.**

## Reaction catalysed

4-phospho-D-erythronate + NAD(+)  $\rightleftharpoons$  (3R)-3-hydroxy-2-oxo-4-phosphonooxybutanoate + NADH

## Comment(s)

- This enzyme catalyzes the second step in the biosynthesis of the coenzyme pyridoxal 5'-phosphate in *Escherichia coli*.
- The reaction occurs predominantly in the reverse direction.
- Other enzymes involved in this pathway are EC 1.2.1.72, EC 2.6.1.52, EC 1.1.1.262, EC 2.6.99.2 and EC 1.4.3.5.

## Cross-references

PROSITE PDOC00063

BRENDA 1.1.1.290

PUMA2 1.1.1.290

PRIAM enzyme-specific profiles 1.1.1.290

Kyoto University LIGAND chemical database 1.1.1.290

IUBMB Enzyme Nomenclature 1.1.1.290

IntEnz 1.1.1.290

MEDLINE Find literature relating to 1.1.1.290

MetaCyc 1.1.1.290

Q8A2E4, PDXB_BACTN;	Q7VRU9, PDXB_BLOFL;	Q83AR8, PDXB_COXBU;
Q8XCR0, PDXB_ECO57;	Q8FFH2, PDXB_ECOL6;	P05459, PDXB_ECOLI;
Q6D2N5, PDXB_ERWCT;	Q7N2B2, PDXB_PHOLL;	Q6LNU2, PDXB_PHOPR;
Q7MV70, PDXB_PORGI;	Q9I3W9, PDXB_PSEAE;	Q88L20, PDXB_PSEPK;



## ENZYME: 1.1.1.290

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**Entry** EC  
1.1.1.290 Enzyme

**Name** 4-phosphoerythronate dehydrogenase;  
PdxB;  
PdxB 4PE dehydrogenase;  
4-O-phosphoerythronate dehydrogenase

**Class** Oxidoreductases  
Acting on the CH-OH group of donors  
With NAD<sup>+</sup> or NADP<sup>+</sup> as acceptor

**Sysname** 4-phospho-D-erythronate:NAD<sup>+</sup> 2-oxidoreductase

**Reaction** 4-phospho-D-erythronate + NAD<sup>+</sup> =  
(3R)-3-hydroxy-2-oxo-4-phosphonooxybutanoate + NADH + H<sup>+</sup>  
[RN:R04210]

**Substrate** 4-phospho-D-erythronate [CPD:C03393];  
NAD<sup>+</sup> [CPD:C00003]

**Product** (3R)-3-hydroxy-2-oxo-4-phosphonooxybutanoate [CPD:C06054];  
NADH [CPD:C00004];  
H<sup>+</sup> [CPD:C00080]

**Comment** This enzyme catalyses the second step in the biosynthesis of the coenzyme pyridoxal 5'-phosphate in *Escherichia coli*. The reaction occurs predominantly in the reverse direction [3]. Other enzymes involved in this pathway are EC 1.2.1.72 (erythrose-4-phosphate dehydrogenase), EC 2.6.1.52 (phosphoserine transaminase), EC 1.1.1.262 (4-hydroxythreonine-4-phosphate dehydrogenase), EC 2.6.99.2 (pyridoxine 5'-phosphate synthase) and EC 1.4.3.5 (pyridoxamine-phosphate oxidase).

**Reference**

- [PMID:2121717]  
Lam HM, Winkler ME.  
Metabolic relationships between pyridoxine (vitamin B6) and serine biosynthesis in *Escherichia coli* K-12.  
*J. Bacteriol.* 172 (1990) 6518-28.
- [PMID:11844765]  
Pease AJ, Roa BR, Luo W, Winkler ME.  
Positive growth rate-dependent regulation of the *pdxA*, *ksgA*, and *pdxB* genes of *Escherichia coli* K-12.  
*J. Bacteriol.* 184 (2002) 1359-69.
- [PMID:8550422]  
Zhao G, Winkler ME.  
A novel alpha-ketoglutarate reductase activity of the *serA*-encoded 3-phosphoglycerate dehydrogenase of *Escherichia coli* K-12 and its possible implications for human 2-hydroxyglutaric aciduria.  
*J. Bacteriol.* 178 (1996) 232-9.
- [PMID:2692566]